



SEQUENCE LISTING

<110> Sebald, Walter

<120> Polypeptide Variants With Increased Heparin-Binding Ability

<130> PA31187-01996/GRI

<140> US 09/913,467

<141> 2000-01-27

<150> DE 199 06 096.7

<151> 1999-02-13

<160> 12

<170> PatentIn Ver. 2.1

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<212> PRT

<213> Artificial sequence

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<222> (1)

<223> K, R or H

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<222> (2)

<223> K, R or H

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<221> MUTAGEN

<222> (3)

<223> K, R, H or no amino acid

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<222> (4)

<223> not K, R, H, but any other amino acid

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<223> not K, R, H, but any other amino acid

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<210> 3
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<223> description artificial sequence:
heparin-binding sequence

<400> 3
Arg Lys Arg Ala
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<210> 4
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<223> description artificial sequence:
heparin-binding sequence

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Met Ala Gln Ala Lys His Lys Gln Arg Lys Arg Ala Arg Lys Arg Leu
1 5 10 15

Lys Ser Ser Cys Lys Arg His Pro Leu Tyr Val Asp Phe Ser Asp Val
20 25 30

Gly Trp Asn Asp Trp Ile Val Ala Pro Pro Gly Tyr His Ala Phe Tyr
35 40 45

Cys His Gly Glu Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr
50 55 60

Asn His Ala Ile Val Gln Thr Leu Val Asn Ser Val Asn Ser Lys Ile
65 70 75 80

Pro Lys Ala Cys Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu
85 90 95

Tyr Leu Asp Glu Asn Glu Lys Val Val Leu Lys Asn Tyr Gln Asp Met
100 105 110

Val Val Glu Gly Cys Gly Cys Arg
115 120

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<212> PRT
<213> Artificial sequence

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<223> description artificial sequence:T4

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Met Ala Gln Ala Lys His Lys Gln Arg Lys Arg Ala Lys His Lys Gln
1 5 10 15

Arg Lys Arg Leu Lys Ser Ser Cys Lys Arg His Pro Leu Tyr Val Asp
20 25 30

Phe Ser Asp Val Gly Trp Asn Asp Trp Ile Val Ala Pro Pro Gly Tyr
35 40 45

His Ala Phe Tyr Cys His Gly Glu Cys Pro Phe Pro Leu Ala Asp His
50 55 60

Leu Asn Ser Thr Asn His Ala Ile Val Gln Thr Leu Val Asn Ser Val
65 70 75 80

Asn Ser Lys Ile Pro Lys Ala Cys Cys Val Pro Thr Glu Leu Ser Ala
85 90 95

Ile Ser Met Leu Tyr Leu Asp Glu Asn Glu Lys Val Val Leu Lys Asn
100 105 110

Tyr Gln Asp Met Val Val Glu Gly Cys Gly Cys Arg
115 120

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<211> 374

<212> DNA

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<223> description artificial sequence:T3
(nucleic acid sequence)

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ctccccggg gtatcacgcc ttttactgcc acggagaatg cccttttcct ctggctgac 180
atctgaactc cactaatcat gccattgttc agacgttggg caactctgtt aactctaaga 240
ttcctaaggc atgctgtgtc ccgacagaac tcagtgttat ctcgatgctg taccttgacg 300
agaatgaaa gggtgtatta aagaactatc aggacatggg tgtggagggt tgtgggtgtc 360
gctagtaagg atcc 374

<210> 8

<211> 386

<212> DNA

<213> Artificial sequence

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(nucleic acid sequence)

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actggattgt gggtcccccg ggggtatcacg ccttttactg ccacggagaa tgcccttttc 180

ctctggctga tcactgaac tccactaatc atgccattgt tcagacgttg gtcaactctg 240
ttaactctaa gattcctaag gcatgctgtg tcccagacaga actcagtgt atctcgatgc 300
tgtaccttga cgagaatgaa aaggttgat taaagaacta tcaggacatg gttgtggagg 360
gttgtgggtg tcgctagtaa ggatcc 386

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<211> 47

<212> DNA

<213> Artificial sequence

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<223> description artificial sequence: artificial

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<210> 10

<211> 47

<212> DNA

<213> Artificial sequence

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<223> description artificial sequence: artificial

<400> 10

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<211> 59

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<210> 12

<211> 59

<212> DNA

<213> Artificial sequence

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<223> description artificial sequence: artificial

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